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Human PCDH2 ASO PC	Human apolipoprote	Human apolipoprote	Human SNAP29 gene	ASO primer #5 to d	Human HTR5A gene a	Tachykinin recepto	'n	<pre>IGF-I oligonucleot</pre>	н	IGF-I oligonucleot	IGFBP3 oligonucleo	3	labeled	eled	Genomic DNA methyl	DNA	Genomic DNA methyl	Genomic DNA methyl	PCR primer #2. Sy	Human FKBP8 allele	PCR primer H-T11A	Tumour antigen ant	ge	HLA sequence 29.	Human ICAM hammerh

ALIGNMENTS

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RESULT 1
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08-JUN-2001; 2001US-296764P.
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                                                                                                                                                                                                                                                                                                                                                   leukocyte
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Claim 1; Page 477; 2038pp; English. Wohlgemuth J, (BIOC-) BIOCARDIA INC Woodward Fry K, Matcuk G, ...
d R, Quertermous T, Altman P, I, Johnson ۲d .. Prentice

(ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring response to treatment in an individual or monitoring response rheumatoid arthritis, osteoarthritis or cytomegalovirus infection.

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Sequence 50 BP; 11 A; 18 C; 6 G; 15 T; 0 other;

Best Loc Matches Query Match

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Similarity

9.3%; llarity 100.0%; Conservative

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Ś 1385 GCCTTCATGCACCTGTCCTTCTAACACGTCGCCTTCAACTGTAATCACA -GCCTTCATGCACCTGTCCTTTCTAACACGTCGCCTTCAACTGTAATCACA DB 1; 0.00018; 0; 50

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ABZ04679
ID ABZC

ABZ04679 standard; DNA; 50 BP

09-JAN-2003 (first entry)

Human leukocyte gene expression profiling probe SEQ ID NO

T7; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe; . 88

Homo sapiens.

WO200257414-A2

25-JUL-2002

22-OCT-2001; 2001WO-US47856

20-OCT-2000; 2000US-241994P. 08-JUN-2001; 2001US-296764P.

New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides

The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 olligonucleotides (ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring response to treatment in an individual. The disease include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure. Systemic lipus erythematosus, the programment of arthering the coarthering of the coarthering rheumatoid arthritis, osteoarthritis or cytomegalovirus

The invention relates to a system for detecting gene expression, who comprises one or two isolated DNA molecules that detect expression gene, where the gene corresponds to any of 8143 oligonucleotides

1; Page 332; 2038pp; English

New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides

WPI; 2002-636525/68.

Ly N, Wohlgemuth J,

woodward R,

K, Matcuk G, Quertermous

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Altman P, Johnson

'TJ Prentice (BIOC-) BIOCARDIA INC

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                                                                                                                                 The present invention relates to human oligonucleotides comprising a single nucleotide polymorphic site (SNP: AAH89219). The present sequence is one such oligonucleotide. The oligonucleotides can be used in forensics, paternity testing, correlation of polymorphisms with phenotypic traits, genetic mapping of phenotypic traits and marker assisted breeding of animals and crop plants.
                                                                                                                                                                                                                        New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly disease, and also in forensics and paternity testing
                                                                                                                  Sequence
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(AFFY-)
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plant breeding;
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AFFYMETRIX INC.
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Variation
                                        DNA sequence analysis; sequencing; protein sequence; protein structure; gene typing; organ donation; bacteria identification; 16s rRNA; HLA; human leukocyte antigen; PCR primer; ss.
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AFFYMETRIX INC.
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Homo sapiens

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Matches 21
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                                                                                                                                                                                                                                                                                                                                    01-NOV-1994
                           Disclosure;
                                                                                                                                            WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 14; Page 49; 66pp; English.
                                                                                                                                                                                       (NITE ) NIPPON TELEGRAPH & TELEPHONE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AMSH ) AMERSHAM PHARMACIA BIOTECH
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les 21; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   restriction
                      Page
                                                                   cDNA and gene expression - digestion with restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Pred. No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis followed
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                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis; gene expression; reverse aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 6; llpp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-1993;
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n with restriction
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                                                                                                                                                                                    C; 1
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                                                                                                                                                                                    G; 18
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                                                                                                            Score
Pred.
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Pred. No.
                                                                                           Mismatches
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                                                                                                               19.4;
No. 63;
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                                                                                                                                                                                    other;
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47;
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RESULT 8
AAQ49436/c
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XX AAQ494
XX Z5-MAR
DT 25-MAR
DT 27-APR
XX Transg
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RESULT 9
AAQ75578/c
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DE Reverse tl
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DE Reverse tl
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DE Reverse tl
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DE Synthetic
XX
Synthetic
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AC AARR-19
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                                                                                                                                                                                                                                                                                                     Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid isolate encoding flavonoid-3'-hydroxylase create transgenic plants with altered petal colour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-1992;
07-JAN-1993;
16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1993-336914/42
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27-APR-1994
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                                                                                                                                                  04-AUG-1995
                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                sequences
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                       01-NOV-1994.
                                              JP06303997-A
                                                                                         aggregate;
                                                                                                   Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                            chain reactions
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                                                                                                                         transcription primer used
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                                                                                                                                                                                              standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plants; altered petal colour; chain reaction; ss.
                                                                                          restriction
                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                            Page 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FLOWER DEV PTY LTD
                                                                                                                                                                                                                                                                                                                                               BP; 1 A;
                                                                                                                                                                                                                                                                                                    Conservative
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(first entry)
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93AU-0006698
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100.0%; Pr
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                                                                                         enzyme;
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                                                                                                                                                                                                                                                                                                                                                                                                                             English.
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                                                                                                                                                                                                                                                                                                             Score 19;
Pred. No.
                                                                                          83
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEO files AQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression
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by digestion with restriction
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